

Supplementary information

Tyrosine Hydroxylase is crucial for maintaining the pupae tanning and immune in *Anopheles sinensis*

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Running title: *AsTH* for *Anopheles sinensis* pupae development

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Figure S1. Alignment analysis of TH amino acid sequences among different insect species. Horizontal line indicates the Biopterin-H domain.

Figure S2. Detection of RNAi off-target effects. a. The pupa cuticle pigmentation pattern at 38th hour of pupation between ds*TH2* and ds*Red* group. Scale bar=500μm. b. The expression level of *AsTH* between ds*TH2* and ds*Red* individuals at 38th hour of pupation. *RPL49* was used as the internal control.

Figure S3. Precise region of cuticle frozen section. The red box region represents the observed sliced section region. Scale bar=100μm.

Figure S4. The expression pattern of *AsTH* gene expression in the egg (eggs were gathered from 1 hour to 1 day after born) stage, larval stage and adult stage, respectively. *RPL49* was used as the internal control.

Figure S5. Spatial expression pattern of *AsTH* in different tissues at 38th hour of pupation. IN represents the integument, FB represents the fat body, (HE+G) represents the mixture of the hemolymph and the gut. The "*" symbol represents treatment with bacteria. *RPL49* was used as the internal control.

Figure S6. The expression levels of 4 representative *Anopheles sinensis* pro-phenoloxidase (PPO) genes (The orthologous genes in *Anopheles gambiae* were expressed in the immune tissues) between ds*TH* and ds*Red* group at 38th hour of pupation. *RPL49* was used as the internal control.

Figure S1

<i>An. gambiae</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>An. coluzzii</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>An. arabiensis</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>An. stephensi</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>An. dirus</i>	: -----MTASVSRENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>An. funestus</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>An. sinesis</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDELPQPKPQEHPVA
<i>Cx. quinquefasciatus</i>	: -MIAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETIVVKQTQTVLDEARAKANEDKIPQEVOQTVEDQNDEEIRMVAVDLPLQPKPEEHVPSA
<i>Ae. aegypti</i>	: -----MEDAYPEQQ-----AVDLPYDPEEIVNSG
<i>Dr. melanogaster</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETLVVKQTQTVLEARSKAN
<i>Ma. sexta</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETLVVKQTQTVSLEEARARANDSGLSEDFI#QDGIGHGNQDNTPTVEDGTQDETQNGLHAD
<i>Bm. mori</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETLVVKQTQTVSLEEARARANDSGLSEDFI#QDGIGHGNQDNTPTVEDGTQDETQNGLHAD
<i>Da. plexippus</i>	: MAVAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETLVVKQTQTVSLEEARARANDSGLSEDFI#QDVSDAAKTEGV---QNECDKNGHLEGG
<i>Tr. castaneum</i>	: -----MIAAAQKNRREMFAIKKSYSIENGVPSSRRSLVDDARFETLVVKQTQTVLDEARVKA-----DIEDKPAQKIDGE aaaqknremfa kksys ElgyPsrrrlsvddarfet vvktqvl ear an
<i>An. gambiae</i>	: DDEKETDAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEAHTGIVVHLESQRSEGVQFDLVVKVDMARANLLQLIRSLRQ
<i>An. coluzzii</i>	: DDEKETDAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEAHTGIVVHLESQRSEGVQFDLVVKVDMARANLLQLIRSLRQ
<i>An. arabiensis</i>	: DDEKETDAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEAHTGIVVHLESQRSEGVQFDLVVKVDMARANLLQLIRSLRQ
<i>An. stephensi</i>	: DDEKEKDAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEAHTGIVVHLESQRSEGVQFDLVVKVDMARANLLQLIRSLRQ
<i>An. dirus</i>	: DDEKDGTAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEAHTGIVVHLESQRSEGVQFDLVVKVDMARANLLQLIRSLRQ
<i>An. funestus</i>	: DDEKEKDAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEAHTGIVVHLESQRSEGVQFDLVVKVDMARANLLQLIRSLRQ
<i>An. sinesis</i>	: NDEDEKTAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEEGYHIVVHLESQRSEGVAFDVLIKVDMARANLLQLIRSLRQ
<i>Cx. quinquefasciatus</i>	: AQDD-ETAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEETHTHIVVHLESQRSEGDGMFLVLIKVDMARANLLQLIRSLRQ
<i>Ae. aegypti</i>	: DEID-----LAGLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEHTHIVVHLESQRSEGVQFDLVIKVDMARANLLQLIRSLRQ
<i>Dr. melanogaster</i>	: -----DYLGTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKAVEHTHIVVHLESQRSEGVQFDLVIKVDMARANLLQLIRSLRQ
<i>Ma. sexta</i>	: DDNAKADEDTLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKTDHDNKGCIQHLETTRPSQVGVDFDALLVKVMSRINLLQLIRSLRQ
<i>Bm. mori</i>	: DDDDADEDTLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKTDHDNKGCVQHLETTRPSQVGVDFDALLVKVMSRINLLQLIRSLRQ
<i>Da. plexippus</i>	: NETGDTDTLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKTDHDNKGCVQHLETTRPSQISGVDFDALLVKVMSRINLLQLIRSLRQ
<i>Tr. castaneum</i>	: -----LSCLTBEEVVLQNAASESPEAEKBVRRAVVVLRQGMGSLCIRILKTDHDNKGCVQHLETTRPNQSDNNLIALVKVMSRINLLQLIRSLRQ D gLTeeE66LqNaasEspeae 6 AA666r64dG6gSL R6Lk 6e G 6 H6E3r s gvqfd L6K6 M R nL6Ql14sLrq3
<i>An. gambiae</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>An. coluzzii</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>An. arabiensis</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>An. stephensi</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>An. dirus</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>An. funestus</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>An. sinesis</i>	: QSFGSVSLLENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMVQ
<i>Cx. quinquefasciatus</i>	: ASFGSVVAALENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMOK
<i>Ae. aegypti</i>	: ASFGSVVAALENNNVVKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADQVYARRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMOK
<i>Dr. melanogaster</i>	: GFSSMNLMAADDNLNWKAPWPFKHASELDNCNHLMLTYEPDLDMNHPGFADKQYVORRKEIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMV
<i>Ma. sexta</i>	: TAAFGVNLMEENNISSTPWFCHASLDLDCNHLMLTYEELDMNHPGFADKEYRERRKQIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMV
<i>Bm. mori</i>	: TAAFGVNLTEBNNISSTPWFCHASLDLDCNHLMLTYEELDMNHPGFADKEYRERRKQIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMV
<i>Da. plexippus</i>	: TSFGAVLMD-NISNKTPWFPHASLDLDCNHLMLTKFEELDMNHPGFADKEYRERRKQIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMV
<i>Tr. castaneum</i>	: SSELHCILIGEDNISNAKNPWFPHASLDLDCNHLMLTYEPDLDMNHPGFADKEYRERRKELIAEIAFAYRYGDPFPIHDYDYTEENKTWAAVFGRVKELMV sf 66 e N6 K PWPF HAS LDNCNHLMLTK5EPdLDMNHPGFAD YR RRK21Ae6FaY4YGDpI Y 3e En TW VF V Im k
<i>An. gambiae</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. coluzzii</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. arabiensis</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. stephensi</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. dirus</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. funestus</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. sinesis</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Cx. quinquefasciatus</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Ae. aegypti</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Dr. melanogaster</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Ma. sexta</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Bm. mori</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Da. plexippus</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>Tr. castaneum</i>	: HACSEYYIAVERKLEDEKIFVKPFRPLPQLEMSDFLRKNTGETLRAAGLITARDFLASLAFRIFQSTQYVRHINSPIHTPEPDICIHELLGHMPILLADPSFA
<i>An. gambiae</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. coluzzii</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. arabiensis</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. stephensi</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. dirus</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. funestus</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. sinesis</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Cx. quinquefasciatus</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Ae. aegypti</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Dr. melanogaster</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Ma. sexta</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Bm. mori</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Da. plexippus</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>Tr. castaneum</i>	: QFSQEIGLASLGADEIEIKLSTVWYFWTVEFLCKEKEDEVKAYGAQLLSAYGELLHAISDKPEIRPFPASTAVQPYQDQEYQPIYYVAESFDDAKEKFR
<i>An. gambiae</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. coluzzii</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. arabiensis</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. stephensi</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. dirus</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. funestus</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. sinesis</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Cx. quinquefasciatus</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Ae. aegypti</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Dr. melanogaster</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Ma. sexta</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Bm. mori</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Da. plexippus</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>Tr. castaneum</i>	: RWVSMMSRPFEVRNPHTFVEVLDSVKEITLVSQNLTEVHLTNIAIKLQPF
<i>An. gambiae</i>	: -----QNTETEGLH13N46 K64 p5

Figure S2

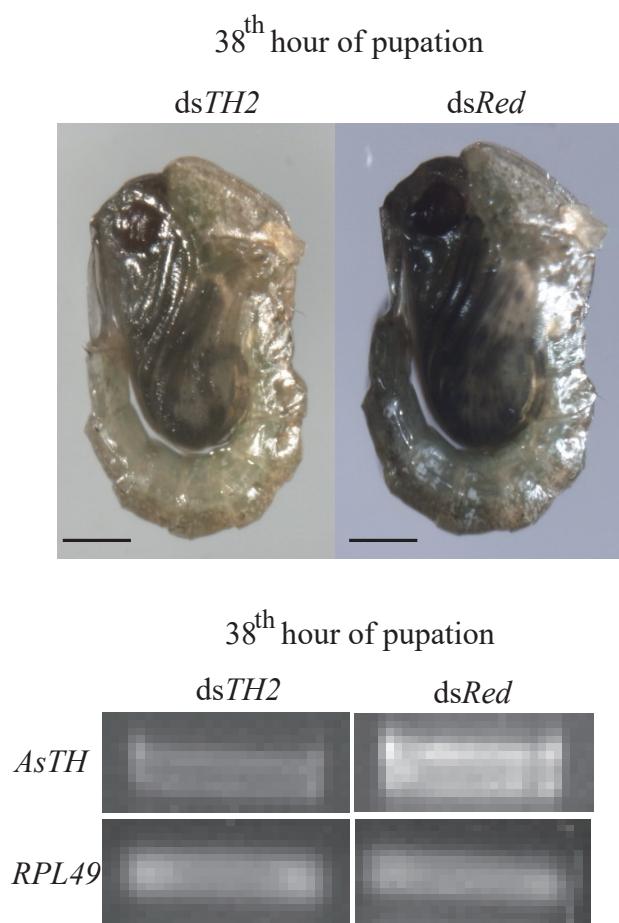


Figure S3

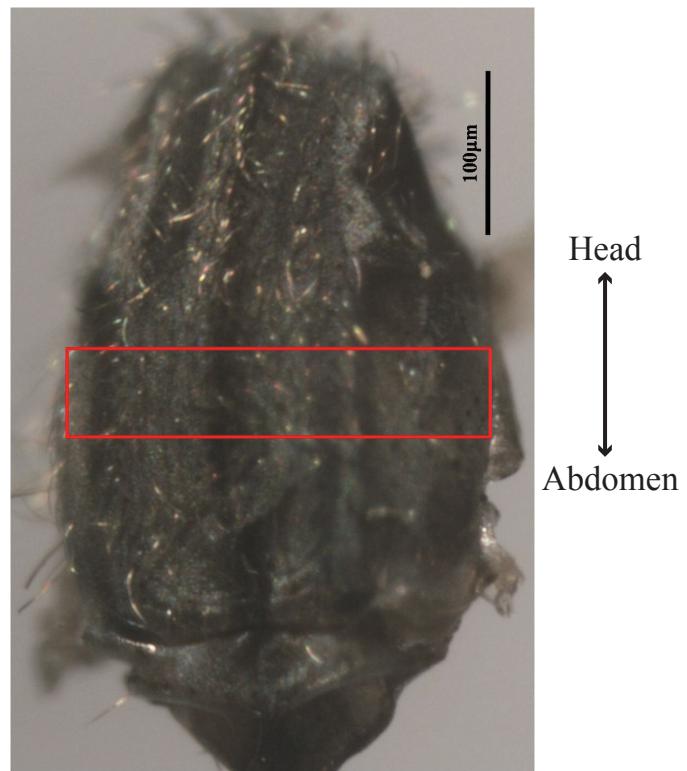


Figure S4

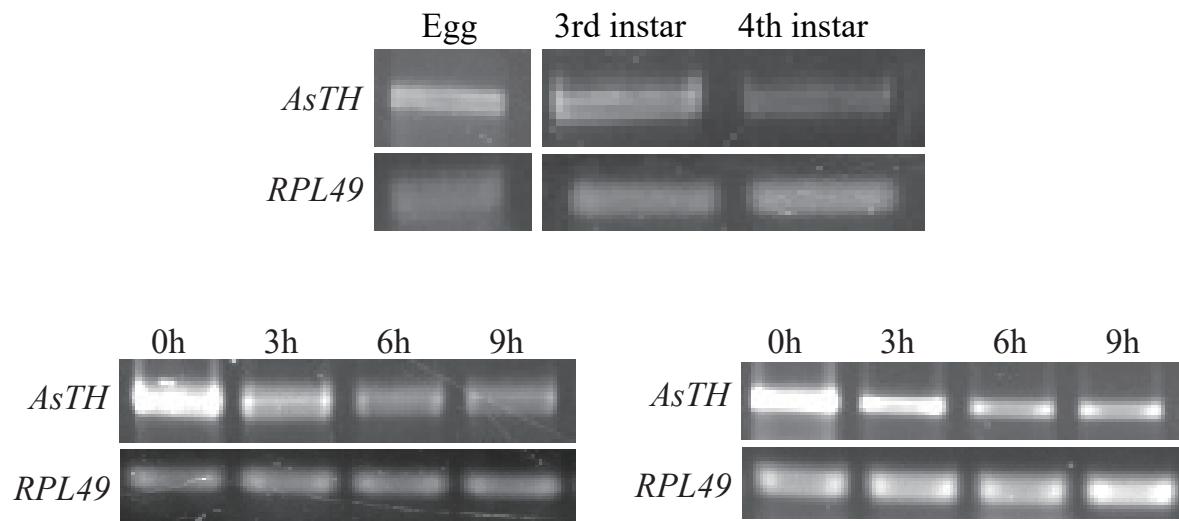


Figure S5

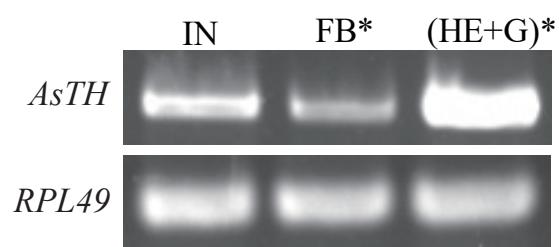


Figure S6

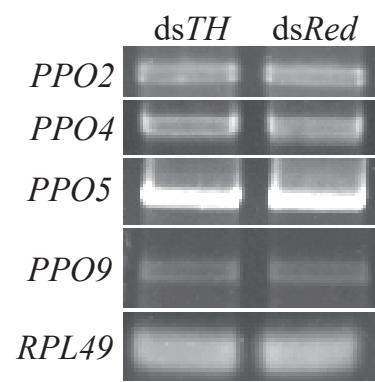


Table S1. Primers used in this study

Primer name	Sequence	
	F primer (5'---3')	R primer (5'---3')
Primers for <i>TH</i> cloning		
<i>AsTH</i> sequence 1	CCGATACCTGCCAAATCCAC	TGAGTCTGACGAGCCCTAACCTCT
<i>AsTH</i> sequence 2	AAGACAAACTCCGCAAGAGGT	CGTGTGGTAGGGCGAGTTG
<i>AsTH</i> sequence 3	CGGAAAGCGGTATGCATTCTATA	TCATCCAGGACGGTTGTTTG
<i>AsTH</i> sequence 4	CCAGCTCAACACCGAAGTGC	AGGGGAAGAGGGAGGATGAAC
3'RACE nested primer	CAGAGTTAGGGCTCGTCAGACTCAT	CGCTACGTAACGGCATGACAGTG
3'RACE primer	CTTATGAGGACGGACAGAGTTAGGG	GCTGTCAACGATACTGCTACGTAACG
Primers used in RT-PCR		
<i>AsTH</i>	GCTGGCGTCCGCATCTT	GCGATGGCATTGGTGAGGT
<i>RPL49</i>	GGAGCCGGTCGGTGTATGT	TTCCCTCTCGGTCGGCTTCG
<i>AsAttacin</i>	CGGGTTTTCTGATTGTGCT	GCGACCGAAGTGCTGCGTAT
<i>AsCecropin-A</i>	ATCTTCTGATTCTGCTGCTGG	GCCTTCTCTGCCGCCTGA
<i>AsCecropin-B</i>	GCCTGTTCGTGGTGTGATG	AGCAGTGCCTCTGAGCCG
<i>AsDefensin</i>	ATCGTGTGGTGTGGTGGTA	GCAGGTTGCCGCTTGAA
<i>AsGambicin</i>	TTCTCCTCGCCGTCCCT	CCGTATCGTGGCAGTCCG
<i>AsPPO 2</i>	CTGGACGAGGAGCGTGGTAT	AGTCACATGGTGCCTAGGATC
<i>AsPPO 4</i>	GTGCGATGGCGGTGTTG	TCGTGGATGTAGCCGAGAATG
<i>AsPPO 5</i>	GTGCCTTCGCCGACATCA	CATTGTATCGCAAACGACCCT
<i>AsPPO 9</i>	CGGGTGTACGCTATGGGA	TCTTCACGGCATCGTCTTCG
Quantitative RT-PCR primers		
<i>AsDDC</i>	GGACAGGGTGGTGGTTATTCA	GATTGCCTTCTCCAGCGTTTC
<i>Aslaccase2</i>	GTTCCCGTAACGAGCACACATT	CTCCAAACCGTCTCGCATC
<i>RPS7</i>	CGGAGAAGATGGCATGGGAGAT	ATAGTGAGCATAGGCCCGGTTA
Primers used for dsRNA template amplification		
<i>AsTH</i> sequence 1	TAATACGACTCACTATAAGGGAGAAAGACAAACTCCGCAAGAGGT	TAATACGACTCACTATAAGGGAGACGTGTGGTAGGGCGAGTTG
<i>AsTH</i> sequence 2	TAATACGACTCACTATAAGGGAGACCGATACTGCCAAATCCAC	TAATACGACTCACTATAAGGGAGATGAGTCTGACGAGCCCTAACCTCT
<i>Red</i>	TAATACGACTCACTATAAGGGAGACTCAAGGTGCGCATGGAG	TAATACGACTCACTATAAGGGAGATGTGGATCTGCCCTTCAG

Table S2. Amino acid identity of AsTH Biopterin_H domain with its orthologs

	<i>An. gambiae iae</i>	<i>An. sine sis</i>	<i>An. dir us</i>	<i>An. funes</i>	<i>An. steph ensi</i>	<i>An. arabie nsis</i>	<i>An. colu zzii</i>	<i>Culex quinquefas ciatus</i>	<i>Aede s aegy pti</i>	<i>Drosoph ila melanog aster</i>	<i>Bom byx mori</i>	<i>Dana us plexip pus</i>	<i>Mand uca sexta</i>	<i>Triboli um castaneum</i>
<i>An. gambiae</i>	100%	96%	99% %	98%	99%	100%	100%	93%	95%	87%	82%	81%	83%	84%
<i>An. sinesis</i>	100% %	96% %	97% %	96%	96%	96%	95%	95%	96%	86%	81%	80%	82%	85%
<i>An. dirus</i>	100% %	98%	99%	99%	99%	99%	93%	95%	87%	87%	83%	81%	83%	85%
<i>An. funestus</i>		100%	98%	98%	98%	98%	95%	95%	87%	83%	81%	81%	83%	85%
<i>An. stephensi</i>			100%	99%	99%	99%	93%	95%	88%	83%	81%	83%	83%	85%
<i>An. arabiensis</i>				100%	100%	100%	93%	95%	87%	82%	81%	83%	83%	84%
<i>An. coluzzii</i>					100%	93%	95%	87%	87%	82%	81%	83%	83%	84%
<i>Culex quinquefas ciatus</i>						100%	95%	85%	81%	79%	80%	83%		
<i>Aedes aegypti</i>							100%	87%	82%	81%	83%	83%	85%	
<i>Drosophila melanogast er</i>								100%	83%	81%	83%	83%	83%	
<i>Bombyx mori</i>									100%	94%	96%	96%	83%	
<i>Danaus plexippus</i>										100%	94%	82%		
<i>Manduca sexta</i>											100%			
<i>Tribolium castaneum</i>												100%		

Table S3. Statistic of RNAi (For Cuticle tanning, at 38th hour of pupation)

Treatment	Injected No.	No. of individuals with tanning impaired (alive)	No. of Death at 38 th hour of pupation
ds <i>TH</i>	58	32	6
ds <i>TH2</i>	42	18	5
dsRed (For ds <i>TH</i>)	49	0	5
dsRed (For ds <i>TH2</i>)	36	0	4